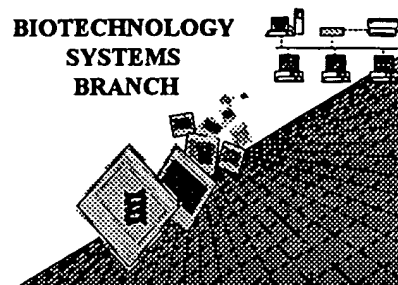


Zaghnoun

# RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY  
SYSTEMS  
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number:

08/894,356A

Art Unit / Team No :

1649

Date Processed by STIC:

1/13/99

**THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.**

**PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:**

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,**
- 2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY**

**THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.**

**IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:**

**ARTI SHAH 703-308-4212**

PAGE: 1

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/894,356ADATE: 01/13/99  
TIME: 16:34:07

INPUT SET: S30733.raw

This Raw Listing contains the General  
Information Section and those Sequences  
containing ERRORS.

## SEQUENCE LISTING

Does Not Comply  
Corrected Diskette Needed

1  
2  
3 (1) General Information:  
4  
5 (i) APPLICANT: ASHIKARI, Toshihiko  
6 TANAKA, Yoshikazu  
7 FUJIWARA, Hiroyuki  
8 NAKAO, Masahiro  
9 FUKUI, Yuko  
10 SAKAKIBARA, Keiko  
11 MIZUTANI, Masako  
12 KUSUMI, Takaaki  
13  
14 (ii) TITLE OF INVENTION: GENE ENCODING A PROTEIN HAVING ACYL  
15 GROUP TRANSFER ACTIVITY  
16  
17 (iii) NUMBER OF SEQUENCES: 31  
18  
19 (iv) CORRESPONDENCE ADDRESS:  
20 (A) ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS, L.L.P.  
21 (B) STREET: 1737 King Street, Suite 500  
22 (C) CITY: Alexandria  
23 (D) STATE: Virginia  
24 (E) COUNTRY: United States  
25 (F) ZIP: 22314-2756  
26  
27 (v) COMPUTER READABLE FORM:  
28 (A) MEDIUM TYPE: Floppy disk  
29 (B) COMPUTER: IBM PC compatible  
30 (C) OPERATING SYSTEM: PC-DOS/MS-DOS  
31 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30  
32  
33 (vi) CURRENT APPLICATION DATA:  
34 (A) APPLICATION NUMBER: US 08/894,356  
35 (B) FILING DATE: 18-AUG-1997  
36 (C) CLASSIFICATION:  
37  
38 (vii) PRIOR APPLICATION DATA:  
39 (A) APPLICATION NUMBER: JP 7-67159  
40 (B) FILING DATE: 17-FEB-1995  
41  
42 (vii) PRIOR APPLICATION DATA:  
43 (A) APPLICATION NUMBER: JP 7-196915  
44 (B) FILING DATE: 29-JUN-1995  
45

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/894,356ADATE: 01/13/99  
TIME: 16:34:08

INPUT SET: S30733.raw

46 (vii) PRIOR APPLICATION DATA:  
47 (A) APPLICATION NUMBER: JP 8-46534  
48 (B) FILING DATE: 30-JAN-1996  
49  
50 (vii) PRIOR APPLICATION DATA:  
51 (A) APPLICATION NUMBER: WO PCT/JP96/00348  
52 (B) FILING DATE: 16-FEB-1996  
53  
54 (viii) ATTORNEY/AGENT INFORMATION:  
55 (A) NAME: Meuth, Donna M.  
56 (B) REGISTRATION NUMBER: 36,607  
57 (C) REFERENCE/DOCKET NUMBER: 001560-308  
58  
59 (ix) TELECOMMUNICATION INFORMATION:  
60 (A) TELEPHONE: (703) 836-6620  
61 (B) TELEFAX: (703) 836-2021  
62  
63

## ERRORED SEQUENCES FOLLOW:

836 (2) INFORMATION FOR SEQ ID NO:6:  
837  
838 (i) SEQUENCE CHARACTERISTICS:  
--> 839 (A) LENGTH: 1521 base pairs  
840 (B) TYPE: nucleic acid  
841 (C) STRANDEDNESS: double  
842 (D) TOPOLOGY: linear  
843  
844 (ii) MOLECULE TYPE: cDNA to mRNA  
845  
846 (iii) HYPOTHETICAL: NO  
847  
848 (iv) ANTI-SENSE: NO  
849  
850 (vi) ORIGINAL SOURCE:  
851 (A) ORGANISM: Lavandula angustifolia  
852 (F) TISSUE TYPE: petal  
853  
854 (vii) IMMEDIATE SOURCE:  
855 (A) LIBRARY: cDNA library  
856 (B) CLONE: pLAT21  
857  
858 (ix) FEATURE:  
859 (A) NAME/KEY: CDS  
860 (B) LOCATION: 1..1352  
861  
862  
863 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:  
864

*1522 shown (next page)*

# RAW SEQUENCE LISTING PATENT APPLICATION US/08/894,356A

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 TIME: 16:34:09

INPUT SET: S30733.raw  
 47

865	NTG ACC ACC CTC CTC GAA TCC TCC CGA GTG GCG CCG CCT CCA GGC ACG	
866	Xaa Thr Thr Leu Leu Glu Ser Ser Arg Val Ala Pro Pro Pro Gly Thr	
867	1 5 10 15	
868		
869	GTG GCT GAG CAG TCA CTC CCG CTC ACC TTC TTC GAC ATG ACG TGG CTG	
870	Val Ala Glu Gln Ser Leu Pro Leu Thr Phe Phe Asp Met Thr Trp Leu	
871	20 25 30	
872		
873	CAT TTC CAC CCC ATG CTT CAG CTT CTC TTC TAC GAA CTC CCC TGT TCC	
874	His Phe His Pro Met Leu Gln Leu Leu Phe Tyr Glu Leu Pro Cys Ser	
875	35 40 45	
876		
877	AAA CCC GCC TTC CTC GAA ACC GTC GTT CCG AAA CTC AAA CAA TCC TTA	
878	Lys Pro Ala Phe Leu Glu Thr Val Val Pro Lys Leu Lys Gln Ser Leu	
879	50 55 60	
880		
881	TCT CTA ACC CTC AAA CAC TTC TTC CCC CTT TCA TGC AAT CTA ATC TAC	
882	Ser Leu Thr Leu Lys His Phe Phe Pro Leu Ser Cys Asn Leu Ile Tyr	
883	65 70 75 80	
884		
885	CCT CTA TCG CCG GAG AAA ATG CCG GAG TTC CGG TAT CAG AAC GGT GAC	
886	Pro Leu Ser Pro Glu Lys Met Pro Glu Phe Arg Tyr Gln Asn Gly Asp	
887	85 90 95	
888		
889	TCG GTT TCT TTC ACG ATT ATG GAG TCT GTC GGA GAT CAT CCG CAT TCC	
890	Ser Val Ser Phe Thr Ile Met Glu Ser Val Gly Asp His Pro His Ser	
891	100 105 110	
892		
893	GCT CAT AAA TAC TAC TGC TTT GCC CCT AGC GAC GAT TAT GAA GAT CTC	
894	Ala His Lys Tyr Tyr Cys Phe Ala Pro Ser Asp Asp Tyr Glu Asp Leu	
895	115 120 125	
896		
897	CAG CTG CCG CCG ATA GTC GAG GAA TCT GAT CGG AAA TTG TTT CAA GTT	
898	Gln Leu Pro Pro Ile Val Glu Glu Ser Asp Arg Lys Leu Phe Gln Val	
899	130 135 140	
900		
901	TTA GCC GTG CAA GTG ACT CTG TTT CCC GGT CGC GGG GTG TGC ATC GGA	
902	Leu Ala Val Gln Val Thr Leu Phe Pro Gly Arg Gly Val Cys Ile Gly	
903	145 150 155 160	
904		
905	ATA ACG ACG CAC CAC ACC GTT AGC GAT GCT CCA TCG TTT GTA GGG TTT	
906	Ile Thr Thr His His Thr Val Ser Asp Ala Pro Ser Phe Val Gly Phe	
907	165 170 175	
908		
909	ATG AAG AGT TGG GCT TCC ATC ACT AAA TTC GGA GGA GAT GAT GAA TTC	
910	Met Lys Ser Trp Ala Ser Ile Thr Lys Phe Gly Gly Asp Asp Glu Phe	
911	180 185 190	
912		
913	TTG GAC GGA AAA GGT GAA TGT TTG CCG GTT TTC GAC CGA TCG CTC GTG	
914	Leu Asp Gly Lys Gly Glu Cys Leu Pro Val Phe Asp Arg Ser Leu Val	
915	195 200 205	
916		
917	AAT TAT CCG CCT AAA TTG GAC ACA TAT TTA TGG AAC AAC GCG CAG AAA	

191

239

287

335

383

431

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527

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623

671

numbers  
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 throughout  
 sequence

# RAW SEQUENCE LISTING PATENT APPLICATION US/08/894,356A

DATE: 01/13/99  
TIME: 16:34:10

INPUT SET: S30733.raw

918	Asn Tyr Pro Pro Lys Leu Asp Thr Tyr Leu Trp Asn Asn Ala Gln Lys	
919	210 215 220	
920		
921	CGT CCG TTG GAA TCG CAG CAT CCA TCT TTA CCG ACG GAT CGG ATT CGA	719
922	Arg Pro Leu Glu Ser Gln His Pro Ser Leu Pro Thr Asp Arg Ile Arg	
923	225 230 235 240	
924		
925	GCT ACC TAC CTT TTC ACC CAA TCT GAA ATT AAG AAA TTG AAG GGT TTG	767
926	Ala Thr Tyr Leu Phe Thr Gln Ser Glu Ile Lys Lys Leu Lys Gly Leu	
927	245 250 255	
928		
929	ATT CAG AGA AAA GCC CCA AAT GTA GTT AAT CTC TCT TCC TTC GTC GCG	815
930	Ile Gln Arg Lys Ala Pro Asn Val Asn Leu Ser Ser Phe Val Ala	
931	260 265 270	
932		
933	ATC GCA GCT TAT ATC TGG ACC GGC ATC GCC AAA TCG GTC GGA GAT TAC	863
934	Ile Ala Ala Tyr Ile Trp Thr Gly Ile Ala Lys Ser Val Gly Asp Tyr	
935	275 280 285	
936		
937	AAA GAC GTG GAT GAC GAC AAA CGC GCT TTC TTT TTA ATT CCG ATC GAT	911
938	Lys Asp Val Asp Asp Asp Lys Arg Ala Phe Phe Leu Ile Pro Ile Asp	
939	290 295 300	
940		
941	TTA AGG CCG CGT TTG GAT CCG CCG GCT CCG GGG AAC TAC TTC GGA AAC	959
942	Leu Arg Pro Arg Leu Asp Pro Pro Ala Pro Gly Asn Tyr Phe Gly Asn	
943	305 310 315 320	
944		
945	TGT CTA TCG TTT GCG ATG GCG AAG ATC CTG CCG CCG GAT TTG GTC GGA	1007
946	Cys Leu Ser Phe Ala Met Ala Lys Ile Leu Arg Arg Asp Leu Val Gly	
947	325 330 335	
948		
949	GAT GAA GGG GTG TTT CGG GCA GCT GAG GCG ATC GCG GCG GAA ATA GAG	1055
950	Asp Glu Gly Val Phe Arg Ala Ala Glu Ala Ile Ala Ala Glu Ile Glu	
951	340 345 350	
952		
953	AAG AGG ACG AGC GAC AAG AAG ATT CTA GAA ACT GTG GAG AAC TGG CCG	1103
954	Lys Arg Thr Ser Asp Lys Lys Ile Leu Glu Thr Val Glu Asn Trp Pro	
955	355 360 365	
956		
957	TCT GAG ATT CGC GAA GCC TTG CAA AAC TGT TAT TTC TCG GTG GCG GGA	1151
958	Ser Glu Ile Arg Glu Ala Leu Gln Asn Cys Tyr Phe Ser Val Ala Gly	
959	370 375 380	
960		
961	TCG AGC AGG CTT GAT CTT TAC GGC GCG GAT TTT GGA TGG GGT AAG GCG	1199
962	Ser Ser Arg Leu Asp Leu Tyr Gly Ala Asp Phe Gly Trp Gly Lys Ala	
963	385 390 395 400	
964		
965	GTG AAG CAA GAG ATA CTG TCG ATT GAT GGA GAG AAG TTT ACG ATG TCG	1247
966	Val Lys Gln Glu Ile Leu Ser Ile Asp Gly Glu Lys Phe Thr Met Ser	
967	405 410 415	
968		
969	TTG TGT AAA CCG AGG GAT GCT GCC GGA GGA TTG GAG GTT GGA TTG TCT	1295
970	Leu Cys Lys Pro Arg Asp Ala Ala Gly Gly L u Glu Val Gly Leu Ser	

↑  
number  
off  
↓

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/894,356ADATE: 01/13/99  
TIME: 16:34:11

INPUT SET: S30733.raw

	420	425	430	
971				
972				
973	TTG CCA AAG GAG GAA TTG CAA GCT TTT GAT GAT TAT TTT GCG GAG GGA			1343
974	Leu Pro Lys Glu Glu Leu Gln Ala Phe Asp Asp Tyr Phe Ala Glu Gly			
975	435	440	445	
976				
977	ATA AAG GGT TGATTAATCA TTTAATCATG TATTATGAAG TTGGATGAAA			1392
978	Ile Lys Gly			
979	450			
980				
981	TCCTCTGTTT CATCTCTATT GTTTAAACAA TAATTTTTTTT CCATTGAACT TTTTGTGAGTC			1452
982				
983	AATAAAAAAAAA AAAAAAAAAA AAAAAAATG AAAAAACTCA GTTATTTTTTT TTTTTTTTTT			1512
984				
985	TTTTTTTTTT			1521
986				
987				

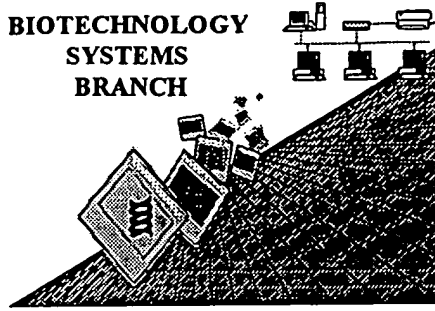
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*Transcribed off**↓*

**SEQUENCE VERIFICATION REPORT**  
**PATENT APPLICATION US/08/894,356A**DATE: 01/13/99  
TIME: 16:34:11**INPUT SET: S30733.raw**

Line	Error	Original Text
839	Entered (1521) and Calc. Seq. Length (1522) differ	(A) LENGTH: 1521 base pairs
865	# of Sequences for line conflicts w/ running total	NTG ACC ACC CTC CTC GAA TCC TCC CGA GTG GC

BIOTECHNOLOGY  
SYSTEMS  
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## Notice of Availability of Checker Program

### Applicant Aid for Biotechnology Computer Readable Form (CRF) Sequence Listing Submissions

The Patent and Trademark Office (PTO) has developed a computer program, called Checker, that will aid applicants in identifying and correcting errors prior to making submissions for compliance with the Requirements for Patent Applications Containing Nucleotide Sequence and/or Amino Acid Sequence Disclosures (Sequence Rules: 37CFR 1.821 through 1.825). Final rules were published in the *Federal Register* (55 FR18230) on May 1, 1990, and in the PTO *Official Gazette* (1114 Off.Gaz.PatOffice 29) on May 15, 1990.

Checker is a DOS-based software program that is intended to assist users in determining whether errors may be present in the sequence listings, and is not intended to guarantee that the submission is error-free.

The most current version of the software is available via computer downloading, details are below. Copies on diskette are also available. Updated software versions will not be automatically mailed out; any updates will be announced in the PTO *Official Gazette*.

The software can be accessed/requested from the following locations:

- 1) Dial-up access through the Internet. Location is <ftp://ftp.uspto.gov>  
The software is in current directory: [pub/checker/](ftp://ftp.uspto.gov/pub/checker/)  
Download all the files. Cost: Free-of-charge
- 3) For diskette copies, mail to: U.S.P.T.O., OEIP, CRYSTAL PARK 3, SUITE 441  
WASHINGTON DC 20231

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*For Further Information, Contact: **Arti Shah at 703-308-4212***